

>AB065623 ACCESSION:AB065623 NID: gi 21928520 dbj AB065623.1 Homo
sapiens gene for seven transmembrane helix receptor,
complete cds, isolate:CBRC7TM_186
Length = 1324

Score = 617 bits (1575), Expect = e-175
Identities = 306/307 (99%), Positives = 307/307 (99%)
Frame = +3

Query: 1 MNHSVVTETFIILGLTKKPELQGIIFLFFLIVYLVAFLGNMLIIIAKIYSNTLHTPMYVFL 60
MNHSVVTETFIILGLTKKPELQGIIFLFFLIVYLVAFLGNMLIIIAKIY+NTLHTPMYVFL
Sbjct: 201 MNHSVVTETFIILGLTKKPELQGIIFLFFLIVYLVAFLGNMLIIIAKIYNNLHTPMYVFL 380

Query: 61 LTLAVVDIICTTSIIPKMLGTMLTSENTISYAGCMSQLFLTWSLGAEMVLFTTMAYDRY 120
LTLAVVDIICTTSIIPKMLGTMLTSENTISYAGCMSQLFLTWSLGAEMVLFTTMAYDRY
Sbjct: 381 LTLAVVDIICTTSIIPKMLGTMLTSENTISYAGCMSQLFLTWSLGAEMVLFTTMAYDRY 560

Query: 121 VAICFPLHYSTIMNHHMCVALLSMVMAIAVTNSWVHTALIMRLTFCGPNTIDHFFCEIPP 180
VAICFPLHYSTIMNHHMCVALLSMVMAIAVTNSWVHTALIMRLTFCGPNTIDHFFCEIPP
Sbjct: 561 VAICFPLHYSTIMNHHMCVALLSMVMAIAVTNSWVHTALIMRLTFCGPNTIDHFFCEIPP 740

Query: 181 LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS 240
LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS
Sbjct: 741 LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS 920

Query: 241 HLTVVTLYYSPVIYTYIRPASSYTFERDKVVAALYTLVTPTLNPMVYSFQNREMQAGIRK 300
HLTVVTLYYSPVIYTYIRPASSYTFERDKVVAALYTLVTPTLNPMVYSFQNREMQAGIRK
Sbjct: 921 HLTVVTLYYSPVIYTYIRPASSYTFERDKVVAALYTLVTPTLNPMVYSFQNREMQAGIRK 1100

Query: 301 VFAFLKH 307
VFAFLKH
Sbjct: 1101VFAFLKH 1121